

Table 1

	Known SAPs		MS/MS from AdML S'some					MS/MS from Ftz S'some					
Reference with link	Protein Name	Mass	Avg XCorr	Total peptides	Unique sequences	Filtered peptides	Filtered unique	Avg XCorr	Total peptide s	Unique sequence s	Filtered peptides	Filtered unique	NCI annotation/GenBank Accession/Other Names

Sm core proteins

sim core proteins													
SW:SMD1_HUMAN	Sm D1	13282	4.081	114	9	83	4	3.977	8	3	6	3	P13641
SW:SMD2_HUMAN	Sm D2	13527	3.897	50	14	26	5	4.012	15	8	10	5	P43330
SW:SMD3_HUMAN	Sm D3	13916	3.859	19	7	10	2	4.169	11	5	4	2	P43331
SW:RUXG_HUMAN	Sm G	8496	2.865	7	2	7	2	3.171	1	1	1	1	Q15357
SW:RSMB_HUMAN	Sm B/B'	24610	3.222	16	9	4	3	3.288	6	5	4	2	P14678
SW:RUXE_HUMAN	Sm E	10804	3.204	5	4	4	3	2.578	4	3	3	2	P08578
SW:RUXF_HUMAN	Sm F	9725	2.799	25	13	3	2	2.731	16	6	4	3	Q15356

U6 snRNA-associated sm-like proteins

OG shRNA-associated sm-time proteins													
SWN:LSM4_HUMAN	lsm4	15350	2.677	11	3	7	2	2.759	7	2	4	2	Q9y4z0
SWN:LSM2_HUMAN	lsm2	10835	4.074	9	6	4	2	3.435	3	2	3	2	Q9y333
SWN:LSM7_HUMAN	lsm7	11602	2.498	3	2	3	2	***					Q9uk45
SWN:LSM6_HUMAN	lsm6	9128	2.989	4	4	1	1	2.338	2	1	2	1	Q9y4y8
SWN:LSM3_HUMAN	lsm3	11714	2.24	10	6	1	1	2.457	6	2	3	1	Q9y4z1
SWN:LSM5_HUMAN	lsm5	9806	***					***					Q9Y4Y9
SWN:LSM8_HUMAN	lsm8	10271	***					***					Q95777

U1 snRNP specific proteins

U1 snRNP specific proteins														
Species	SW:RU17_HUMAN	U1 70k	70081	3.291	14	6	6	2	3.246	3	3	1	1	P08621
	SW:RU1C_HUMAN	U1 C	17394	3.771	2	1	2	1	3.853	2	2	1	1	P09234
	SW:RU1A_HUMAN	U1 A	31280	2.208	7	5	1	1	2.775	7	5	2	2	P09012

U2 snRNP specific proteins

GP:AF054284	1	SAP155	145815	3.475	124	43	64	23	2.743	52	26	29	15	
SW:S145	HUMAN	SAP145	97657	3.228	67	33	30	12	3.084	33	20	15	8	Q13435
SW:S114	HUMAN	SAP 114	88886	3.549	57	30	29	14	3.042	37	22	19	11	Q15459
GP:AJ001443	1	SAP 130	135592	3.471	16	10	9	5	3.096	9	6	7	5	
PIR2:A55749		SAP61	58849	3.308	33	21	20	11	2.921	15	7	7	3	
SW:SP62	HUMAN	SAP 62	49196	2.992	12	7	8	4	2.24	7	6	1	1	Q15428
SW:SP49	HUMAN	SAP49	44386	3.543	7	4	4	2	4.207	5	5	2	2	Q15427
SW:RU2A_HUMAN		U2 A'	28444	3.25	39	19	19	9	2.863	18	11	12	8	P09661
SW:RU2B_HUMAN		U2 B"	25486	4.024	11	6	6	1	5.28	6	3	1	1	P08579
AAK94041		d14	14584	***					***					

U5 snRNP specific proteins

O14834 - specific proteins														
SWN:U520_HUMAN	U5-200	194479	3.395	150	62	101	36	2.929	154	56	96	32	lost link, DEIH-box	
GP-AB007510	U5-220	237875	0.307	111	42	59	25	2.778	143	53	77	29	lost link	
GNP-BC002360	U5-116	109478	3.308	70	38	47	22	2.812	68	34	45	16	EF-2-like	
GNP-BC001666	U5-102	106925	0.046	6	5	4	4	2.744	9	6	3	3	hPrp6, hPrp1, SPF107	
GNP-BC002366	U5-100	95583	2.763	27	21	17	12	2.474	22	14	13	10	DEAD-box	
GP-AF090988	U5-40	39299	3.865	4	3	3	2	3.169	6	3	5	3		
SW:DM1_HUMAN	U5-15	16786	3.649	3	3	1	1	2.247	1	1	1	1	O14834, dm1 homolog	

U4/U6,U5 snRNP specific proteins

PIR2:T50839	U4/U6-90k	77529	3.335	15	13	9	8	2.558	11	8	6	5	hPrp3
GPN:BC007424_1	U4/U6-60k	58321	3.037	10	8	6	5	2.852	10	7	6	4	hPrp4
GP:AF083385_1	SPF30	26711	3.46	13	7	7	3	2.836	5	3	4	2	
PIR2:T00034	Trn-snRNP 110k	90255	3.147	22	16	11	11	2.918	19	15	11	9	SPF90/SART1/hSnu66
GP:AF353989_1	Trn-snRNP 65k	65415	2.46	2	2	1	1	***					hSad1p
SW:NHPX_HUMAN	Trn-snRNP15 5kD	14174	2.926	7	4	5	3	2.861	2	2	1	1	P55769, AF155235, hSnu13p
PIR1:S64705	CyP-60	58823	3.089	12	9	11	8	3.194	1	1	1	1	cyclophilin-like protein CyP-60
GPN:AF271652_1	PPIL3b	18155	3.162	9	6	5	4	3.185	5	5	2	2	cyclophilin-like protein PPIL3b (PPIL3)

Step II proteins

Step 1: Proteins													
SW-PR16_HUMAN	hPrp16	140473	3.054	7	6	6	5	2.569	5	3	3	2	Q92620, DEAH-box
GP-AF038392_1	hPrp17	65521	3.2	10	6	6	4	2.585	3	2	1	1	
GPB-BC010634_1	hSlu7	68343	3.448	6	6	2	2	2.636	4	4	2	2	
GPB-BC000794_1	hPrp18	39860	4.918	1	1	1	1	***					
SW-DDX8_HUMAN	hPrp22	139315	3.364	12	9	10	7	2.71	15	13	9	8	Q14562, HRH1, DEAH-box

mRNP proteins

SWN:RBM8_HUMAN	Y14	19889	2.842	7	5	5	3	2.461	5	4	3	2	Q9y5s9, RBM8
AF047002	Alv	26861	3.32	8	6	2	1	2.841	9	6	3	2	NM_005782, BEF
SW:MGN_HUMAN	meqoh	17164	2.934	7	5	3	2	2.713	7	4	5	3	P50606
PIR2:JC4525	RNPS1	34208	4.132	7	5	3	2	3.191	4	3	3	2	RNA-binding protein E5 1

SR proteins

SW:SFR1_HUMAN	SF2/ASF	27613	2.828	27	14	19	10	2.852	42	12	36	9	Q07955
SW:SFR3_HUMAN	SRp20	19330	2.762	16	4	11	3	2.764	11	4	8	3	P23152
SW:SFR7_HUMAN	9G8	27367	2.529	11	4	7	3	2.416	6	3	4	3	Q16629
SW:SFR6_HUMAN	SRp55	39568	3.181	6	4	4	3	3.027	8	4	6	3	Q13247
SW:SFR4_HUMAN	SRp75	56792	2.381	5	4	3	2	2.537	6	3	2	2	Q08170
SW:SFR5_HUMAN	SRp40	31264	3.327	2	2	2	2	3.018	4	1	4	1	Q13243
SW:SFR9_HUMAN	SRp30c	25542	2.759	5	3	2	2	2.414	10	6	5	4	Q13242
SW:SFR2_HUMAN	SC35	25575	2.974	2	2	1	1	***					Q01130
GP:AF048977_1	SRm160	93519	3.559	5	4	2	2	3.578	7	5	5	3	
SW:SFRB_HUMAN	p54	53542	3.633	9	5	5	3	2.276	2	2	1	1	Q05519, SFRS11
SW:SFR8_HUMAN	SWAP	104821	***					2.282	4	4	1	1	Q12872, SFRS8

Other dead box/helicase proteins

SW:HE47_HUMAN	UAP56	48991	3.131	7	7	3	3	2.54	7	5	5	3	hBAT1, p47, HE47, Q13838
SW:DD15_HUMAN	hPrp43	92829	3.237	14	11	12	9	2.778	11	10	5	4	O43143
SW:DD17_HUMAN		72371	2.894	13	10	11	8	2.554	20	11	14	6	Q92841, p72
SW:DDX3_HUMAN		73243	3.233	6	5	6	5	2.54	3	3	2	2	O00571, nlp2
SW:DDX5_HUMAN		69148	2.643	10	7	6	6	2.515	9	6	5	3	P17844, p68, mann
SW:DD16_HUMAN	hPrp2	119172	2.64	10	7	6	5	2.613	9	6	5	3	O60231
SW:DDX9_HUMAN		140877	3.063	7	6	2	1	2.75	9	7	16	5	Q08211, ndh ii
GP:AF106680_1	hPrp5	117266	2.539	2	2	1	1	3.455	1	1	1	1	RNA helicase

Other SAPs

SW:CB80_HUMAN	CBP 80	91839	2.846	28	16	20	12	2.672	38	18	20	10	Q09161
SW:CB20_HUMAN	CBP20	18001	3.281	8	4	2	1	2.482	8	5	4	3	P52298
SW:U2AF_HUMAN	U2 AF65	53501	3.641	15	8	11	5	3.758	6	5	1	1	P26368
SW:U2AG_HUMAN	U2AF35	27872	2.77	3	3	2	2	3.942	2	2	2	2	Q01081
GPN:AY029347_1	hPrp4/kinase	116973	3.838	6	3	5	2	2.967	3	3	2	2	serine/threonine-protein kinase (PRP4)
GPN:BC008719_1	hPrp19	55181	3.165	43	17	19	9	2.931	20	10	12	7	nuclear matrix protein NMP200, WD repeat
GP:AL050389_1	hPrp31	55424	3.049	8	6	4	3	3.207	4	4	3	3	
GP:AF049523_1	hFBP 11		2.917	8	4	7	3	2.864	11	5	6	3	huntingtin-interacting protein HYBA/EBP11
GP:AF255443_1	hCRN	99201	3.032	25	16	16	11	3.318	8	6	4	4	CGI-201, TPR-repeat
Y08765/Y08766	SF1/mBBP	68632	***					***					several isoforms

SAPs with other function

PIR2:T08599	CA150	123960	2.925	56	28	33	15	2.684	32	23	21	13	
SW:SKIP_HUMAN	skip	61494	3.507	22	18	8	7	2.896	18	9	12	5	snw1, nuclear receptor coactivator ncoa-62
GPN:BC007871_1	SPF45	44962	2	2	1	1		3.118	1	1	1	1	G-patch, RRM, DNA damage repair
GP:AF083383_1	SPF38	34290	2.876	4	3	2	2	3.823	5	4	4	3	WD

Proteins enriched in H complex

SW:PTB_HUMAN	hnRNP I/PTB	57221	4.13	25	18	11	6	2.206	3	3	1	1	P26599
SW:ROA1_HUMAN	hnRNP A1	38715	3.48	15	11	8	6	3.354	20	13	8	5	P09651
SW:ROA2_HUMAN	hnRNP A2/B1	37430	3.515	3	3	2	2	2.653	14	8	8	4	P22626
SWN:ROR_HUMAN	hnRNP R	70943	3.074	10	9	7	7	3.006	15	9	11	6	O43390
SW:ROC_HUMAN	hnRNP C1/C2	33299	2.828	7	6	6	6	2.791	34	10	27	10	P07910
SW:ROK_HUMAN	hnRNP K	50976	3.482	5	5	5	5	3.356	16	9	9	6	Q07244
SW:ROA0_HUMAN	hnRNP A0	30841	4.284	7	3	4	2	4.373	3	2	3	2	Q13151
SW:ROH1_HUMAN	hnRNP H	49229	3.47	6	5	4	3	***					P31943
SW:ROL_HUMAN	hnRNP L	60187	2.884	6	6	3	3	2.989	55	21	37	14	P14866
SWN:ROD_HUMAN	hnRNP D0	38434	2.879	4	3	3	2	3.365	8	5	4	1	Q14103
SW:ROM_HUMAN	hnRNP M	77469	2.815	5	5	3	3	***					P52272
GPN:BC001616_1	hnRNP A/B	30586	3.667	2	1	2	1	3.06	1	1	1	1	
SW:ROA3_HUMAN	hnRNP A3	39686	3.655	6	5	2	2	2.823	7	6	3	2	P51991
SW:ROF_HUMAN	hnRNP F	45672	3.429	2	2	2	2	***					P52597
SW:ROG_HUMAN	hnRNP G	42404	***					3.036	10	4	7	3	P38159
PIR2:B54857	NF-AT 90k	73339	3.117	7	4	7	4	2.919	9	8	7	6	
PIR2:A54857	NF-AT 45k	44697	3.436	7	6	4	4	3.197	8	7	6	5	
GP:AF037448_1	Gry-rbp	69633	3.882	3	3	1	1	3.149	9	5	6	4	RRM RNA binding protein
GPN:BC008875_1	PUF60	58171	3.316	57	29	24	13	3.217	29	18	5	4	siah binding protein 1, FBP interacting repressor, PTB

Table 2

Reference with link	NCI annotation	Mass	MS/MS from AdML S'some	Avg XCorr	Total peptides	Unique sequences	Filtered peptides	Filtered unique	Avg XCorr	Total peptides	Unique sequences	Filtered peptides	Filtered unique
Potentially new SAPs													
GP:AF356524_1	nuclear receptor transcription cofactor (SHARP)	402248	3.39	32	25	16	13	2.829	23	18	10	8	
GPN:BC007208_1	HCNP, XPA-binding protein 2	100010	3.241	27	18	17	12	2.779	9	8	6	5	
GP:AC004858_3	U1 small nonnucleoprotein 1SNRP homolog, Pole(A) binding protein	94122	3.331	50	26	33	15	3.454	24	15	13	8	
PIR2:155595	splicing factor	58657	3.346	32	14	18	7	2.9	10	5	4	2	
GP:AB034205_1	cisplatin resistance-associated overexpressed protein	51466	3.45	14	9	10	6	3.149	8	7	7	6	
SW:BUB3_HUMAN	mitotic checkpoint protein bub3	37155	2.982	6	5	6	5	2.956	7	4	5	3	
SWN:ELV1_HUMAN	elav-like protein 1 (hu-antigen r) (hur)	36062	2.787	2	2	1	1	2.93	10	6	5	3	
GPN:BC001621_1	Npw38-binding protein NpwBP	69998	2.64	11	9	7	5	2.332	5	4	2	1	
SW:RED_HUMAN	red protein (rer protein) (ik factor) (cytokine ik)	65630	2.781	9	7	5	4	2.517	4	4	2	2	
SW:Z07_HUMAN	O43670, zinc finger protein 207	50751	3.199	10	6	2	2	2.64	6	5	1	1	
SW:YB1_HUMAN	y box binding protein-1 (yb-1) (ccaaat-binding transcription factor 1 subunit a) (cbf-a) enhancer factor	35924	4.636	10	9	1	1	4.316	6	5	2	1	
GPN:BC003376_1	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)	36092	3.878	2	1	1	1	3.862	5	1	3	1	
PIR2:A53545	nuclear matrix protein p84	75627	3.557	13	10	6	5	2.99	29	13	15	7	
GP:AF155096_1	NY-REN-6 antigen, partial cds		2.973	17	6	4	2	2.82	11	4	6	3	
SW:IF4N_HUMAN	eukaryotic initiation factor 4a-like nuk-34	46833	3.147	14	12	9	9	2.383	11	8	6	5	
SWN:CRK7_HUMAN	cell division cycle 2-related protein kinase 7 (cdc2-related protein kinase 7) (crkrs)	164155	3.146	11	8	6	3	3.07	7	6	5	4	
GPN:BC001403_1	pre-mRNA cleavage factor Im (25kD)	26227	3.246	11	7	7	5	3.511	6	5	4	3	
GP:AF044333_1	pleiotropic regulator 1 (PLRG1)	57194	3.768	13	10	7	6	3.656	4	4	3	3	
SW:G10_HUMAN	P41223, human homolog of xenopus maternal g10 protein (edg-2)	16844	2.277	2	2	1	1	2.431	4	2	3	1	
GP:AJ271745_1	double-stranded RNA binding nuclear protein DRBP76 (ILF3 gene)	76033	2.804	2	1	1	1	2.41	5	3	3	1	
GP:AJ276706_1	partial mRNA for WTAP (wilm's tumor associating protein) (hFL2D)		3.572	2	1	1	1	3.715	2	2	2	2	
GP:AJ279080_1	putative transcription factor (ORF1), ORF1 S1 RNA binding protein	104804	3.544	3	3	3	3	3.033	3	3	2	2	
SW:GCFC_HUMAN	gc-rich sequence dna-binding factor homolog, Q9Y6b6	29010	3.167	3	3	3	3	2.655	2	1	2	1	
GP:U70667_1	Fas-ligand associated factor 1 (FLAF1), WWP/WW motif		2.51	10	6	3	1	2.616	5	4	2	2	
GPN:BC005152_1	similar to mouse Gli3 or D. melanogaster transcription factor IIB	22774	3.154	1	1	1	1	2.606	2	2	2	2	
SWN:R856_HUMAN	Q92804, tata-binding protein associated factor 2n (ma-binding protein 56) (tbf168)	61830	2.726	3	2	3	2	2.011	3	2	2	1	
SWN:CYCK_HUMAN	O75909, cyclin k	41293	3.653	2	1	1	1	4.355	2	2	1	1	
GPN:BC003015_1	Similar to expressed sequence 2 embryonic lethal		3.363	3	3	3	3	4.199	1	1	1	1	
GP:AB016088_1	RNA binding protein, partial cds		3.764	10	5	5	2	4.054	3	3	1	1	
SW:SP18_HUMAN	O00422, sin3 associated polypeptide p18	17561	3.414	2	2	1	1	3.435	5	5	1	1	
GPN:BC002548_1	Similar to Moloney leukemia virus 10	113671	3.215	4	4	2	2	3.434	2	2	1	1	
PIR2:I38191	nucleic acid binding protein (fragment)		4.65	1	1	1	1	3.364	2	2	1	1	
SW:CIRP_HUMAN	cold-inducible ma-binding protein (glycine-rich ma-binding protein grp)	18648	3.037	5	3	4	2	3.349	2	2	1	1	
GP:AF112222_1	nuclear protein SDK3	81584	3.026	4	3	1	1	3.311	5	3	1	1	
GP:L76159_1	Facioscapulohumeral muscular dystrophy region gene-1	29172	3.363	2	1	2	1	3.079	1	1	1	1	
SW:GR78_HUMAN	78 kda glucose-regulated protein precursor (grp 78) (immunoglobulin heavy chain binding protein) (bip)	72116	4.29	3	2	1	1	2.955	1	1	1	1	
GPN:BC000495_1	CD2 antigen (cytoplasmic tail)-binding protein 2	37646	3.374	4	3	2	2	2.887	4	4	1	1	
SW:MFA1_HUMAN	P55081, microfilament-associated protein 1	51855	2.687	6	5	3	3	2.777	3	3	1	1	
GP:AF015044_1	EH-binding protein, binds EH domains of eps15		3.132	6	6	3	3	2.771	1	1	1	1	
GPN:BC010381_1	nuclear matrix protein p84	75666	2.555	5	5	3	3	2.768	6	4	1	1	
GP:AF361746_1	endothelial cell-selective adhesion molecule (ESAM), immunoglobulin superfamily, contains V and C2 domains	41208	3.24	3	1	3	1	2.555	1	1	1	1	
GP:AJ271741_1	partial ILF3 gene for interleukin enhancer binding factor 3	25972	2.303	3	2	2	1	2.379	4	3	1	1	
SW:EF11_HUMAN	elongation factor 1-alpha 1 (ef-1-alpha-1) (elongation factor tu) (ef-tu)	50141	2.852	6	5	5	4	2.377	1	1	1	1	
SW:DBPA_HUMAN	dna-binding protein a (cold shock domain protein a) (single-strand dna binding protein nf-gmb)	40060	3.194	10	8	5	3	2.374	2	2	1	1	
Hypothetical new SAPs													
PIR2:T00365_??	hypothetical protein KIAA0670 (fragment)		3.207	77	37	44	18	3.001	72	33	45	17	
PIR2:T02345	hypothetical protein KIAA0324 (fragment), SRm300, matrix related		3.069	58	30	31	17	2.885	37	25	20	13	
SW:Y017_HUMAN	hypothetical protein kaa0017	44606	3.668	39	20	24	9	3.192	19	11	13	5	
GP:AB007892_1	KIAA0432 mRNA	58171	3.417	51	32	23	16	2.962	27	21	15	12	
PIR2:T00333	hypothetical protein KIAA0560	163966	3.292	31	21	18	12	2.643	30	19	17	12	
PIR2:T12455	hypothetical protein DKFZp564H2023.1 (fragments)		2.898	35	18	20	10	2.931	67	21	40	10	
SW:YS64_HUMAN	hypothetical protein s164 (fragment)		3.614	12	6	7	4	2.998	16	7	12	6	

GP:AK023659_1	clone PLACE1009798, weakly similar to RLR1 PROTEIN	73795	2.902	12	8	11	7	2.698	33	16	25	12
PIR2:T46386	hypothetical protein DKFZp434P011.1 (fragment)		3.071	15	11	11	8	2.602	17	12	12	9
GN:BC003118_1	clone MGC-2655 IMAGE 3537243, mRNA, complete cds	34849	3.259	5	3	4	2	2.565	14	6	11	4
GP:AF151059_1	HSPC225 (FBP11 related)	24297	2.938	14	6	8	5	2.66	15	5	10	4
PIR2:I39463	gene anonymous protein - human	78536	3.161	5	4	3	2	2.629	16	12	8	6
SWN:CG80_HUMAN	hypothetical protein cgi-110 (protein hspc175)	14585	3.8	13	7	7	2	3.082	5	4	4	3
GN:BC006397_1	Similar to hypothetical protein FLJ12479, clone MGC:13150 IMAGE:4298786	102135	3.052	9	5	4	2	2.882	7	6	4	3
GP:AL031668_1	Human DNA sequence from clone RP1-64K7 on chromosome 20q11.21-11.23 Contains the EIF2S2 gene for eukaryotic translation initiation factor 2 subunit 2 (beta, 38kD), a putative novel gene, the gene for heterogeneous nuclear ribonucleoprotein RALY or aut	32214	2.707	2	2	1	1	2.547	5	3	4	2
GP:D38552_1	KIAA0073 gene, partial cds, The ha1539 protein is related to cyclophilin		2.924	14	10	9	7	2.585	8	6	4	4
GP:AB046824_1	Homo sapiens mRNA for KIAA1604 protein, partial cds, Start codon is not identified		3.596	16	11	11	8	3.345	8	6	3	3
GP:AF130096_1	FLC0586 PRO2855 mRNA, complete cds	34655	3.323	10	4	8	3	3.174	3	2	3	2
GN:BC003048_1	CGI-124 protein	18237	3.038	8	4	7	4	2.716	4	3	3	2
GP:AB018344_1	Homo sapiens mRNA for KIAA0801 protein, complete cds	117461	3.37	24	14	14	9	2.586	10	9	3	2
GP:AK027098_1	Homo sapiens cDNA: FLJ23445 fis, clone HSI01721, unnamed protein product	23671	2.498	4	3	3	2	2.566	10	6	3	3
GN:BC003402_1	hypothetical protein FLJ10290, clone MGC:4943 IMAGE:3449258, mRNA, complete cds	46896	3.376	8	8	3	3	2.5	5	4	3	2
PIR2:T12485	hypothetical protein DKFZp564O2082.1 - human	28722	4.171	4	4	1	1	3.224	6	4	2	2
GN:BC004442_1	Similar to RIKEN cDNA 5830446M03 gene, clone MGC 4036 IMAGE:2820683, mRNA, complete	32992	2.902	4	4	3	3	3	2	1	2	1
PIR2:T17232	hypothetical protein DKFZp434I116.1 - human (fragment)		3.077	4	4	2	2	2.956	4	4	2	2
GN:BC006474_1	clone IMAGE 2820942, mRNA, partial cds		3.342	2	2	2	2	2.813	4	4	2	2
GN:BC004122_1	Similar to RIKEN cDNA 5830446M03 gene, clone MGC:11203 IMAGE 3927759, mRNA	10870	3.599	2	2	1	1	2.707	2	1	2	1
SW:Y105 HUMAN	hypothetical protein kaa0105	17801	3.049	3	3	3	3	2.563	2	1	2	1
GN:BC002876_1	hypothetical protein FLJ10805	57544	3.093	12	6	8	4	2.561	3	2	2	1
GP:AF161497_1	Homo sapiens HSPC148 mRNA, complete cds	26610	2.734	5	3	2	1	2.403	5	3	2	2
PIR2:T12531	hypothetical protein DKFZp434B194.1	96820	3.006	10	10	5	5	2.336	7	6	2	2
GP:AB023146_1	KIAA0929 protein, partial cds		3.205	22	16	15	11	2.015	9	8	2	2
GN:BC003359_1	hypothetical protein, clone MGC 5267 IMAGE 2900332, mRNA, complete cds	33855	2.687	2	2	1	1	4.451	1	1	1	1
GN:BC000198_1	Similar to CG11985 gene product, clone MGC:3133 IMAGE 3352960, mRNA, complete cds	10135	3.837	33	2	4	1	4.05	2	1	1	1
PIR2:T02672	hypothetical protein R31449_3 - human (fragment)		3.456	6	6	3	3	3.604	4	4	1	1
PIR2:T46935	hypothetical protein DKFZp434D199.1	14263	3.763	6	6	4	4	3.106	1	1	1	1
GP:AF161433_1	HSPC315 mRNA, partial cds		3.043	3	2	2	1	2.955	2	2	1	1
GP:AF132955_1	CGI-21 protein mRNA	37542	2.706	4	2	2	1	2.795	1	1	1	1
GP:AK000741_1	cDNA FLJ20734 fis, clone HEP08523, unnamed protein product	70516	3.219	3	3	1	1	2.692	2	2	1	1
SW:Y052 HUMAN	hypothetical protein kaa0052 (fragment)		2.916	6	6	3	3	2.611	3	3	1	1
GP:AB011132_1	KIAA0560 protein, partial cds		2.525	1	1	1	1	2.595	1	1	1	1
GN:BC004258_1	Homo sapiens, hypothetical protein PRO1741, clone MGC:10753 IMAGE:3347345, mRNA, complete cds	65691	3.813	3	2	2	2	2.567	1	1	1	1
GN:BC006350_1	clone MGC:13125 IMAGE:4111572	70521	2.77	9	9	7	7	2.419	1	1	1	1
GP:AL512685_1	cDNA DKFZp547K202 (from clone DKFZp547K202); WD-protein The frame shift was determined manually	75399	3.222	3	3	1	1	2.373	4	4	1	1
GP:AL023804_1	Human DNA sequence from clone RP4-633O20 on chromosome 20q11.23-12 Contains 5' end of a gene similar to Bos taurus P14 protein (P14L) ESTs, CA repeat(D20S859), STSs and GSSs, complete sequence, Also similar to Drosophila CG11964 protein	32895	2.81	4	4	3	3	2.321	2	2	1	1